



Form PTO 1449 US Department of Commerce Patent and Trademark Office	ATTY DOCKET NO: P-IS 4988	SERIAL NO. 09/993,312
	APPLICANT: Ideker and Hood	
INFORMATION DISCLOSURE STATEMENT BY APPLICANT	FILING DATE: November 13, 2001	GROUP: 2736

TECHNOLOGY CENTER AREA  
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U.S. PATENT DOCUMENTS

EXAM. INITIALS	DOCUMENT NUMBER	DATE	NAME	CLASS	SUB- CLASS	FILING DATE

FOREIGN PATENT DOCUMENTS

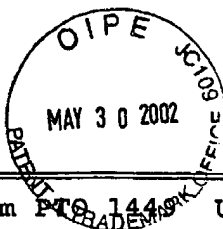
EXAM. INITIALS	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUB- CLASS	TRANSLATION (YES/NO)
CS	WO 00/45322	08/03/00	<del>PCT</del> WIP0	G06F	19/00	

OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages)

CS	Cheng and Church, "Biclustering of Expression Data," <u>Proc. Int. Conf. Intell. Syst. Mol. Biol.</u> 8:93-103 (2000)
	Eisenberg et al., "Protein function in the post-genomic era," <u>Nature</u> 405:823-826 (2000)
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	Hughes et al., "Functional discovery via a Compendium of Expression Profiles," <u>Cell</u> 102:109-126 (2000)
	Ideker, T., "Statement of Research Interests" (Fall 2000)
	Kuffner et al., "Pathway analysis in metabolic data bases via differential metabolic display (DMD)," <u>Bioinformatics</u> 16:825-836 (2000)
↓	Lockhart & Winzeler, "Genomics, gene expression and DNA arrays," <u>Nature</u> 405:827-836 (2000)

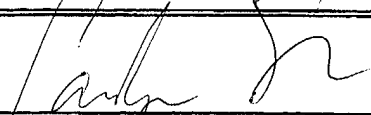
EXAMINER <i>Barry S.</i>	DATE CONSIDERED 12/4/01
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CS		Marcotte et al., "A combined algorithm for genome-wide prediction of protein function," <u>Nature</u> 402:83-86 (1999)
		Proux et al., "A pragmatic information extraction strategy for gathering data on genetic interactions," <u>Proc. Int. Conf. Intell. Syst. Mol. Biol.</u> 8:279-285 (2000)
		Schwikowski et al., "A network of protein - protein interactions in yeast," <u>Nature Biotechnol.</u> 18:1257-1261 (2000)
		Tohsato et al., "A multiple alignment algorithm for metabolic pathway analysis using enzyme hierarchy," <u>Proc. Int. Conf. Intell. Syst. Mol. Biol.</u> 8:376-383 (2000)
		van Someren et al., "Linear modeling of genetic networks from experimental data," <u>Proc. Int. Conf. Intell. Syst. Mol. Biol.</u> 8:355-366 (2000)
		Yona and Levitt, "Towards a complete map of the protein space based on a unified sequence and structure analysis of all known proteins," <u>Proc. Int. Conf. Intell. Syst. Mol. Biol.</u> 8:395-406 (2000)
		Zien et al., "Analysis of Gene Expression Data with pathway scores," <u>Proc. Int. Conf. Intell. Syst. Mol. Biol.</u> 8:407-417 (2000)
		Timecourse Data: Metabolic Chart, <a href="http://cmqm.stanford.edu/pbrown/explore/metachart.html">http://cmqm.stanford.edu/pbrown/explore/metachart.html</a> ; pp. 1-2 (2000)
		Product Description: "GeneSpring Lite," Silicone Genetics Website, <a href="http://www.sigmetrics.com/Products/GeneSpring/Overlays.htm">http://www.sigmetrics.com/Products/GeneSpring/Overlays.htm</a> updated as of Feb. 29, 2000

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